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P.5

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RAW SEQUENCE LISTING DATE: 10/17/2001
PATENT APPLICATION: US/09/825,561A TIME: 12:23:45

Input Set : A:\00-22SEQ.txt
Output Set: N:\CRF3\10172001\I825561A.raw

ENTERED

```
      4 <110> APPLICANT: Sprecher, Cindy A.
      5       Novak, Julia E.
      6       West, James W.
      7       Presnell, Scott R.
      8       Holly, Richard D.
      9       Nelson, Andrew J.
     12 <120> TITLE OF INVENTION: SOLUBLE ZALPHA11 CYTOKINE RECEPTORS
     15 <130> FILE REFERENCE: 00-22
C--> 17 <140> CURRENT APPLICATION NUMBER: US/09/825,561A
C--> 17 <141> CURRENT FILING DATE: 2000-04-05
     17 <150> PRIOR APPLICATION NUMBER: US 60/194,731
     18 <151> PRIOR FILING DATE: 2000-04-05
     20 <150> PRIOR APPLICATION NUMBER: US 60/222,121
     21 <151> PRIOR FILING DATE: 2000-07-28
     23 <160> NUMBER OF SEQ ID NOS: 86
     25 <170> SOFTWARE: FastSEQ for Windows Version 3.0
     27 <210> SEQ ID NO: 1
     28 <211> LENGTH: 1614
     29 <212> TYPE: DNA
     30 <213> ORGANISM: Homo sapiens
     32 <220> FEATURE:
     33 <221> NAME/KEY: CDS
     34 <222> LOCATION: (1)...(1614)
     36 <400> SEQUENCE: 1
     37  atg ccg cgt ggc tgg gcc gcc ccc ttg ctc ctg ctg ctg ctc cag gga       48
     38  Met Pro Arg Gly Trp Ala Ala Pro Leu Leu Leu Leu Leu Leu Gln Gly
     39   1               5                  10                  15
     41  ggc tgg ggc tgc ccc gac ctc gtc tgc tac acc gat tac ctc cag acg       96
     42  Gly Trp Gly Cys Pro Asp Leu Val Cys Tyr Thr Asp Tyr Leu Gln Thr
     43              20                  25                  30
     45  gtc atc tgc atc ctg gaa atg tgg aac ctc cac ccc agc acg ctc acc      144
     46  Val Ile Cys Ile Leu Glu Met Trp Asn Leu His Pro Ser Thr Leu Thr
     47          35                  40                  45
     49  ctt acc tgg caa gac cag tat gaa gag ctg aag gac gag gcc acc tcc      192
     50  Leu Thr Trp Gln Asp Gln Tyr Glu Glu Leu Lys Asp Glu Ala Thr Ser
     51      50                  55                  60
     53  tgc agc ctc cac agg tcg gcc cac aat gcc acg cat gcc acc tac acc      240
     54  Cys Ser Leu His Arg Ser Ala His Asn Ala Thr His Ala Thr Tyr Thr
     55  65                  70                  75                  80
     57  tgc cac atg gat gta ttc cac ttc atg gcc gac gac att ttc agt gtc      288
     58  Cys His Met Asp Val Phe His Phe Met Ala Asp Asp Ile Phe Ser Val
     59                  85                  90                  95
     61  aac atc aca gac cag tct ggc aac tac tcc cag gag tgt ggc agc ttt      336
     62  Asn Ile Thr Asp Gln Ser Gly Asn Tyr Ser Gln Glu Cys Gly Ser Phe
     63              100                 105                 110
     65  ctc ctg gct gag agc atc aag ccg gct ccc cct ttc aac gtg act gtg      384
     66  Leu Leu Ala Glu Ser Ile Lys Pro Ala Pro Pro Phe Asn Val Thr Val
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RAW SEQUENCE LISTING DATE: 10/17/2001
PATENT APPLICATION: US/09/825,561A TIME: 12:23:45

Input Set : A:\00-22SEQ.txt
Output Set: N:\CRF3\10172001\I825561A.raw

```
67          115                 120                 125
69  acc ttc tca gga cag tat aat atc tcc tgg cgc tca gat tac gaa gac      432
70  Thr Phe Ser Gly Gln Tyr Asn Ile Ser Trp Arg Ser Asp Tyr Glu Asp
71      130                 135                 140
73  cct gcc ttc tac atg ctg aag ggc aag ctt cag tat gag ctg cag tac      480
74  Pro Ala Phe Tyr Met Leu Lys Gly Lys Leu Gln Tyr Glu Leu Gln Tyr
75  145                 150                 155                 160
77  agg aac cgg gga gac ccc tgg gct gtg agt ccg agg aga aag ctg atc      528
78  Arg Asn Arg Gly Asp Pro Trp Ala Val Ser Pro Arg Arg Lys Leu Ile
79                  165                 170                 175
81  tca gtg gac tca aga agt gtc tcc ctc ctc ccc ctg gag ttc cgc aaa      576
82  Ser Val Asp Ser Arg Ser Val Ser Leu Leu Pro Leu Glu Phe Arg Lys
83              180                 185                 190
87  gac tcg agc tat gag ctg cag gtg cgg gca ggg ccc atg cct ggc tcc      624
88  Asp Ser Ser Tyr Glu Leu Gln Val Arg Ala Gly Pro Met Pro Gly Ser
89          195                 200                 205
91  tcc tac cag ggg acc tgg agt gaa tgg agt gac ccg gtc atc ttt cag      672
92  Ser Tyr Gln Gly Thr Trp Ser Glu Trp Ser Asp Pro Val Ile Phe Gln
93      210                 215                 220
95  acc cag tca gag gag tta aag gaa ggc tgg aac cct cac ctg ctg ctt      720
96  Thr Gln Ser Glu Glu Leu Lys Glu Gly Trp Asn Pro His Leu Leu Leu
97  225                 230                 235                 240
99  ctc ctc ctg ctt gtc ata gtc ttc att cct gcc ttc tgg agc ctg aag      768
100  Leu Leu Leu Leu Val Ile Val Phe Ile Pro Ala Phe Trp Ser Leu Lys
101                 245                 250                 255
103  acc cat cca ttg tgg agg cta tgg aag aag ata tgg gcc gtc ccc agc      816
104  Thr His Pro Leu Trp Arg Leu Trp Lys Lys Ile Trp Ala Val Pro Ser
105              260                 265                 270
107  cct gag cgg ttc ttc atg ccc ctg tac aag ggc tgc agc gga gac ttc      864
108  Pro Glu Arg Phe Phe Met Pro Leu Tyr Lys Gly Cys Ser Gly Asp Phe
109          275                 280                 285
111  aag aaa tgg gtg ggt gca ccc ttc act ggc tcc agc ctg gag ctg gga      912
112  Lys Lys Trp Val Gly Ala Pro Phe Thr Gly Ser Ser Leu Glu Leu Gly
113      290                 295                 300
115  ccc tgg agc cca gag gtg ccc tcc acc ctg gag gtg tac agc tgc cac      960
116  Pro Trp Ser Pro Glu Val Pro Ser Thr Leu Glu Val Tyr Ser Cys His
117  305                 310                 315                 320
119  cca cca cgg agc ccg gcc aag agg ctg cag ctc acg gag cta caa gaa     1008
120  Pro Pro Arg Ser Pro Ala Lys Arg Leu Gln Leu Thr Glu Leu Gln Glu
121                 325                 330                 335
123  cca gca gag ctg gtg gag tct gac ggt gtg ccc aag ccc agc ttc tgg     1056
124  Pro Ala Glu Leu Val Glu Ser Asp Gly Val Pro Lys Pro Ser Phe Trp
125              340                 345                 350
127  ccg aca gcc cag aac tcg ggg ggc tca gct tac agt gag gag agg gat     1104
128  Pro Thr Ala Gln Asn Ser Gly Gly Ser Ala Tyr Ser Glu Glu Arg Asp
129          355                 360                 365
131  cgg cca tac ggc ctg gtg tcc att gac aca gtg act gtg cta gat gca     1152
132  Arg Pro Tyr Gly Leu Val Ser Ile Asp Thr Val Thr Val Leu Asp Ala
133      370                 375                 380
```

RAW SEQUENCE LISTING DATE: 10/17/2001
PATENT APPLICATION: US/09/825,561A TIME: 12:23:45

Input Set : A:\00-22SEQ.txt
Output Set: N:\CRF3\10172001\I825561A.raw

```
135  gag ggg cca tgc acc tgg ccc tgc agc tgt gag gat gac ggc tac cca     1200
136  Glu Gly Pro Cys Thr Trp Pro Cys Ser Cys Glu Asp Asp Gly Tyr Pro
137  385                 390                 395                 400
139  gcc ctg gac ctg gat gct ggc ctg gag ccc agc cca ggc cta gag gac     1248
140  Ala Leu Asp Leu Asp Ala Gly Leu Glu Pro Ser Pro Gly Leu Glu Asp
141                  405                 410                 415
143  cca ctc ttg gat gca ggg acc aca gtc ctg tcc tgt ggc tgt gtc tca     1296
144  Pro Leu Leu Asp Ala Gly Thr Thr Val Leu Ser Cys Gly Cys Val Ser
145              420                 425                 430
147  gct ggc agc cct ggg cta gga ggg ccc ctg gga agc ctc ctg gac aga     1344
148  Ala Gly Ser Pro Gly Leu Gly Gly Pro Leu Gly Ser Leu Leu Asp Arg
149          435                 440                 445
151  cta aag cca ccc ctt gca gat ggg gag gac tgg gct ggg gga ctg ccc     1392
152  Leu Lys Pro Pro Leu Ala Asp Gly Glu Asp Trp Ala Gly Gly Leu Pro
153      450                 455                 460
155  tgg ggt ggc cgg tca cct gga ggg gtc tca gag agt gag gcg ggc tca     1440
156  Trp Gly Gly Arg Ser Pro Gly Gly Val Ser Glu Ser Glu Ala Gly Ser
157  465                 470                 475                 480
159  ccc ctg gcc ggc ctg gat atg gac acg ttt gac agt ggc ttt gtg ggc     1488
160  Pro Leu Ala Gly Leu Asp Met Asp Thr Phe Asp Ser Gly Phe Val Gly
161                  485                 490                 495
163  tct gac tgc agc agc cct gtg gag tgt gac ttc acc agc ccc ggg gac     1536
164  Ser Asp Cys Ser Ser Pro Val Glu Cys Asp Phe Thr Ser Pro Gly Asp
165              500                 505                 510
167  gaa gga ccc ccc cgg agc tac ctc cgc cag tgg gtg gtc att cct ccg     1584
168  Glu Gly Pro Pro Arg Ser Tyr Leu Arg Gln Trp Val Val Ile Pro Pro
169          515                 520                 525
173  cca ctt tcg agc cct gga ccc cag gcc agc                           1614
174  Pro Leu Ser Ser Pro Gly Pro Gln Ala Ser
175      530                 535
178 <210> SEQ ID NO: 2
179 <211> LENGTH: 538
180 <212> TYPE: PRT
181 <213> ORGANISM: Homo sapiens
183 <400> SEQUENCE: 2
184  Met Pro Arg Gly Trp Ala Ala Pro Leu Leu Leu Leu Leu Leu Gln Gly
185   1               5                  10                  15
186  Gly Trp Gly Cys Pro Asp Leu Val Cys Tyr Thr Asp Tyr Leu Gln Thr
187              20                  25                  30
188  Val Ile Cys Ile Leu Glu Met Trp Asn Leu His Pro Ser Thr Leu Thr
189          35                  40                  45
190  Leu Thr Trp Gln Asp Gln Tyr Glu Glu Leu Lys Asp Glu Ala Thr Ser
191      50                  55                  60
192  Cys Ser Leu His Arg Ser Ala His Asn Ala Thr His Ala Thr Tyr Thr
193  65                  70                  75                  80
194  Cys His Met Asp Val Phe His Phe Met Ala Asp Asp Ile Phe Ser Val
195                  85                  90                  95
196  Asn Ile Thr Asp Gln Ser Gly Asn Tyr Ser Gln Glu Cys Gly Ser Phe
197              100                 105                 110
```

RAW SEQUENCE LISTING DATE: 10/17/2001
PATENT APPLICATION: US/09/825,561A TIME: 12:23:45

Input Set : A:\00-22SEQ.txt
Output Set: N:\CRF3\10172001\I825561A.raw

```
198  Leu Leu Ala Glu Ser Ile Lys Pro Ala Pro Pro Phe Asn Val Thr Val
199          115                 120                 125
200  Thr Phe Ser Gly Gln Tyr Asn Ile Ser Trp Arg Ser Asp Tyr Glu Asp
201      130                 135                 140
202  Pro Ala Phe Tyr Met Leu Lys Gly Lys Leu Gln Tyr Glu Leu Gln Tyr
203  145                 150                 155                 160
204  Arg Asn Arg Gly Asp Pro Trp Ala Val Ser Pro Arg Arg Lys Leu Ile
205                  165                 170                 175
206  Ser Val Asp Ser Arg Ser Val Ser Leu Leu Pro Leu Glu Phe Arg Lys
207              180                 185                 190
208  Asp Ser Ser Tyr Glu Leu Gln Val Arg Ala Gly Pro Met Pro Gly Ser
209          195                 200                 205
210  Ser Tyr Gln Gly Thr Trp Ser Glu Trp Ser Asp Pro Val Ile Phe Gln
211      210                 215                 220
212  Thr Gln Ser Glu Glu Leu Lys Glu Gly Trp Asn Pro His Leu Leu Leu
213  225                 230                 235                 240
214  Leu Leu Leu Leu Val Ile Val Phe Ile Pro Ala Phe Trp Ser Leu Lys
215                  245                 250                 255
216  Thr His Pro Leu Trp Arg Leu Trp Lys Lys Ile Trp Ala Val Pro Ser
217              260                 265                 270
218  Pro Glu Arg Phe Phe Met Pro Leu Tyr Lys Gly Cys Ser Gly Asp Phe
219          275                 280                 285
220  Lys Lys Trp Val Gly Ala Pro Phe Thr Gly Ser Ser Leu Glu Leu Gly
221      290                 295                 300
222  Pro Trp Ser Pro Glu Val Pro Ser Thr Leu Glu Val Tyr Ser Cys His
223  305                 310                 315                 320
224  Pro Pro Arg Ser Pro Ala Lys Arg Leu Gln Leu Thr Glu Leu Gln Glu
225                  325                 330                 335
226  Pro Ala Glu Leu Val Glu Ser Asp Gly Val Pro Lys Pro Ser Phe Trp
227              340                 345                 350
228  Pro Thr Ala Gln Asn Ser Gly Gly Ser Ala Tyr Ser Glu Glu Arg Asp
229          355                 360                 365
230  Arg Pro Tyr Gly Leu Val Ser Ile Asp Thr Val Thr Val Leu Asp Ala
231      370                 375                 380
232  Glu Gly Pro Cys Thr Trp Pro Cys Ser Cys Glu Asp Asp Gly Tyr Pro
233  385                 390                 395                 400
234  Ala Leu Asp Leu Asp Ala Gly Leu Glu Pro Ser Pro Gly Leu Glu Asp
235                  405                 410                 415
236  Pro Leu Leu Asp Ala Gly Thr Thr Val Leu Ser Cys Gly Cys Val Ser
237              420                 425                 430
238  Ala Gly Ser Pro Gly Leu Gly Gly Pro Leu Gly Ser Leu Leu Asp Arg
239          435                 440                 445
240  Leu Lys Pro Pro Leu Ala Asp Gly Glu Asp Trp Ala Gly Gly Leu Pro
241      450                 455                 460
242  Trp Gly Gly Arg Ser Pro Gly Gly Val Ser Glu Ser Glu Ala Gly Ser
243  465                 470                 475                 480
244  Pro Leu Ala Gly Leu Asp Met Asp Thr Phe Asp Ser Gly Phe Val Gly
245                  485                 490                 495
246  Ser Asp Cys Ser Ser Pro Val Glu Cys Asp Phe Thr Ser Pro Gly Asp
```

RAW SEQUENCE LISTING DATE: 10/17/2001
PATENT APPLICATION: US/09/825,561A TIME: 12:23:45

Input Set : A:\00-22SEQ.txt
Output Set: N:\CRF3\10172001\I825561A.raw

```
247              500                 505                 510
248  Glu Gly Pro Pro Arg Ser Tyr Leu Arg Gln Trp Val Val Ile Pro Pro
249          515                 520                 525
250  Pro Leu Ser Ser Pro Gly Pro Gln Ala Ser
251      530                 535
253 <210> SEQ ID NO: 3
254 <211> LENGTH: 696
255 <212> TYPE: DNA
256 <213> ORGANISM: Homo sapiens
259 <220> FEATURE:
260 <221> NAME/KEY: CDS
261 <222> LOCATION: (1)...(696)
263 <400> SEQUENCE: 3
264  ctg aac acg aca att ctg acg ccc aat ggg aat gaa gac acc aca gct       48
265  Leu Asn Thr Thr Ile Leu Thr Pro Asn Gly Asn Glu Asp Thr Thr Ala
266   1               5                  10                  15
268  gat ttc ttc ctg acc act atg ccc act gac tcc ctc agt gtt tcc act       96
269  Asp Phe Phe Leu Thr Thr Met Pro Thr Asp Ser Leu Ser Val Ser Thr
270                20                  25                  30
272  ctg ccc ctc cca gag gtt cag tgt ttt gtg ttc aat gtc gag tac atg      144
273  Leu Pro Leu Pro Glu Val Gln Cys Phe Val Phe Asn Val Glu Tyr Met
274            35                  40                  45
276  aat tgc act tgg aac agc agc tct gag ccc cag cct acc aac ctc act      192
277  Asn Cys Thr Trp Asn Ser Ser Ser Glu Pro Gln Pro Thr Asn Leu Thr
278        50                  55                  60
280  ctg cat tat tgg tac aag aac tcg gat aat gat aaa gtc cag aag tgc      240
281  Leu His Tyr Trp Tyr Lys Asn Ser Asp Asn Asp Lys Val Gln Lys Cys
282   65                  70                  75                  80
284  agc cac tat cta ttc tct gaa gaa atc act tct ggc tgt cag ttg caa      288
285  Ser His Tyr Leu Phe Ser Glu Glu Ile Thr Ser Gly Cys Gln Leu Gln
286                  85                  90                  95
288  aaa aag gag atc cac ctc tac caa aca ttt gtt gtt cag ctc cag gac      336
289  Lys Lys Glu Ile His Leu Tyr Gln Thr Phe Val Val Gln Leu Gln Asp
290              100                 105                 110
292  cca cgg gaa ccc agg aga cag gcc aca cag atg cta aaa ctg cag aat      384
293  Pro Arg Glu Pro Arg Arg Gln Ala Thr Gln Met Leu Lys Leu Gln Asn
294          115                 120                 125
296  ctg gtg atc ccc tgg gct cca gag aac cta aca ctt cac aaa ctg agt      432
297  Leu Val Ile Pro Trp Ala Pro Glu Asn Leu Thr Leu His Lys Leu Ser
298      130                 135                 140
302  gaa tcc cag cta gaa ctg aac tgg aac aac aga ttc ttg aac cac tgt      480
303  Glu Ser Gln Leu Glu Leu Asn Trp Asn Asn Arg Phe Leu Asn His Cys
304  145                 150                 155                 160
306  ttg gag cac ttg gtg cag tac cgg act gac tgg gac cac agc tgg act      528
307  Leu Glu His Leu Val Gln Tyr Arg Thr Asp Trp Asp His Ser Trp Thr
308                  165                 170                 175
310  gaa caa tca gtg gat tat aga cat aag ttc tcc ttg cct agt gtg gat      576
311  Glu Gln Ser Val Asp Tyr Arg His Lys Phe Ser Leu Pro Ser Val Asp
312              180                 185                 190
```

Use of n and / or Xaa has been detected in the Sequence Listing. Review the Sequence Listing to ensure a corresponding explanation is present in the <220> to <223> fields of each sequence using n or Xaa.

VERIFICATION SUMMARY DATE: 10/17/2001
PATENT APPLICATION: **US/09/825,561A** TIME: 12:23:46

Input Set : **A:\00-22SEQ.txt**
Output Set: **N:\CRF3\10172001\I825561A.raw**

L:17 M:270 C: Current Application Number differs, Replaced Current Application No
L:17 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:484 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:485 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:486 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:487 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:488 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:489 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:490 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:491 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:492 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:493 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:494 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:510 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:511 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:512 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:513 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:514 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:515 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:516 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:517 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:518 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:519 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:520 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:521 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:850 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:2392 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:2393 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:2394 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:2395 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:2396 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:2397 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:2398 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:2399 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:2400 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:2401 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:2402 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:2403 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:2404 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:2419 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:67
L:2578 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71
L:2581 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71